Amendments to the Claims:

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

1-30.	(cancelled)
1 50.	(Carrochica)

31.	Plant c	ells, comprising heterologous DNA encoding an EG307 polypeptide,
wherein said p	olypep	tide is capable of increasing the yield of a plant, wherein said polypeptide
selected from	the gro	up consisting of:
	a)	a polypeptide encoded by a polynucleotide selected from the group
consisting of S	SEQ ID	NO:4, SEQ ID NO:5, SEQ ID NO: 91, SEQ ID NO:33, SEQ ID NO:34,
and SEQ ID N	<u>10:35 ;</u>	
	b)	a polypeptide encoded by a polynucleotide having at least 84% sequence
identity to a p	olynucl	eotide in a);
	c)	a polypeptide comprising SEQ ID NO:6 or SEQ ID NO:36; and
	d)	a polypeptide having at least 84% sequence identity to a polypeptide of c)
32.	A prop	pagation material of a transgenic plant comprising the transgenic plant cell
according to c		
33.	A tran	sgenic plant containing heterologous DNA which encodes an EG307
polypeptide th	nat is ex	pressed in plant tissue, wherein said polypeptide increases the yield of the
plant, and saic	d polype	eptide selected from the group consisting of:
	a)	a polypeptide encoded by a polynucleotide selected from the group
consisting of S	SEQ ID	NO:4, SEQ ID NO:5, SEQ ID NO: 91, SEQ ID NO:33, SEQ ID NO:34,
and SEQ ID N	NO:35;	
	b)	a polypeptide encoded by a polynucleotide having at least 84% sequence
identity to a p	olynucl	eotide in a);
	c)	a polypeptide comprising SEQ ID NO:6 or SEQ ID NO:36; and
	d)	a polypeptide having at least 84% sequence identity to a polypeptide of c)
and which cor	nfere cui	ostantially the same yield as the polypentide of c)

34.	An isolated polynucleotide which includes a promoter operably linked to a
polynucleotic	de that encodes the an EG307 gene in plant tissue, said polynucleotide selected from
the group cor	nsisting of:
	a) a polynucleotide selected from the group consisting of SEQ ID NO:4,
SEQ ID NO:	5, SEQ ID NO: 91, SEQ ID NO:33, SEQ ID NO:34, and SEQ ID NO:35;
	b) a polynucleotide having at least 84% sequence identity to a polynucleotide
<u>of a),</u>	
	c) a polynucleotide encoding a polypeptide comprising SEQ ID NO: 6; and
	d) a polynucleotide encoding a polypeptide comprising a protein having at
least 84% sec	quence identity to SEQ ID NO: 6, and which confers substantially the same yield as
the polypepti	de of c).
35.	The inelated neluminal action of Claim 24, subarain acid neluminal action is a
	The isolated polynucleotide of Claim 34, wherein said polynucleotide is a
recombinant	polynucleotide.
36.	(currently amended) The method polynucleotide of claim 34, wherein the
promoter is th	ne promoter native to an EG307 gene.
37-44	. (cancelled).
45.	A transfected host cell comprising a host cell transfected with a construct
comprising a	promoter, enhancer or intron polynucleotide from an evolutionarily significant
EG307 polyn	ucleotide or any combination thereof, operably linked to a polynucleotide encoding
a reporter pro	tein, wherein said EG307 polynucleotide is selected from the group consisting of:
	a) a polynucleotide comprising selected from the group consisting of SEQ ID
NO:4, SEQ II	D NO:5, SEQ ID NO: 91, SEQ ID NO:33, SEQ ID NO:34, and SEQ ID NO:35;
	b) a polynucleotide having at least 84% sequence identity to a polynucleotide
of a),	
	c) a polynucleotide encoding a polypeptide comprising SEQ ID NO: 6; and
	d) a polynucleotide encoding a polypeptide comprising a protein having at
least 84% seq	uence identity to SEQ ID NO: 6, and which confers substantially the same yield as
the polypeptic	de of c).

46	б. А г	nethod of identifying an agent which may modulate yield, said method
comprisir	ng contact	ting at least one candidate agent with a plant or cell comprising an EG307
gene, who	erein the	agent is identified by its ability to modulate yield, and wherein said EG307
gene com	prises a p	polynucleotide selected from the group consisting of:
	a)	a polynucleotide selected from the group consisting of SEQ ID NO:4,
SEQ ID 1	NO:5, SE	Q ID NO: 91, SEQ ID NO:33, SEQ ID NO:34, and SEQ ID NO:35;
	b)	a polynucleotide having at least 84% sequence identity to a polynucleotide
<u>of a),</u>		
	c)	a polynucleotide encoding a polypeptide comprising SEQ ID NO: 6; and
	<u>d)</u>	a polynucleotide encoding a polypeptide comprising a protein having at
least 84%	sequence	e identity to SEQ ID NO: 6, and which confers substantially the same yield as
the polyp	eptide of	<u>c)</u> .
4.5		
47		method of Claim 46, wherein the plant or cell is transfected with a
polynucle	otide <u>of</u>	a), b), c), or d) encoding and EG307 gene.
48	3. (car	ncelled)
49	. The	method of claim 46, wherein said identified agent modulates yield by
modulatir	ng a funct	ion of the polynucleotide encoding the polypeptide.
50). The	method of claim 46, wherein said identified agent modulates yield by
modulating a function of the polypeptide.		
51	l. (cancel	lled)
52	2. (cancel	led)
53	3. (curren	atly amended) A method of producing an EG307 polypeptide comprising:
a)	providing	g a cell transfected with a polynucleotide encoding an EG307 polypeptide
positione	d for expi	ression in the cell;
b) culturing the transfected cell under conditions for expressing the polynucleotide; and		
c)	isolating	the EG307 polypeptide, wherein said polypeptide is selected from the group

consisting of:

i)	a polypeptide encoded by a polynucleotide selected from the group
consisting of SEQ	ID NO:4, SEQ ID NO:5, SEQ ID NO: 91, SEQ ID NO:33, SEQ ID NO:34,
and SEQ ID NO:3	<u>5;</u>
ii)	a polypeptide encoded by a polynucleotide having at least 84% sequence
identity to a polyn	ucleotide in i);
iii)	a polypeptide comprising SEQ ID NO:6 or SEQ ID NO:36; and
iv)	a polypeptide having at least 84% sequence identity to a polypeptide of
<u>iii</u>).	

54-78. (cancelled)

79. (new) A method of detecting a yield-increasing gene in a plant cell comprising contacting the EG307 gene or a portion thereof greater than 12 nucleotides length with a preparation of nucleic acids from the plant cell under hybridization conditions providing detection of nucleic acid molecule sequences having about 50% or greater sequence identity to the a nucleic acid molecule selected from the group consisting of SEQ ID NO:4, SEQ ID NO:5, SEQ ID NO:91, SEQ ID NO:33, SEQ ID NO:34, and SEQ ID NO:35.